



Sequence Listing

#5

<110> Gosdard, Audrey
Woodowski, Paul J.
Gurney, Austin L.
Watanabe, Colin K.
Wood, William I.

<120> NOVEL POLYPEPTIDES HAVING SEQUENCE SIMILARITY TO
CYTOKINE RECEPTORS AND NUCLEIC ACIDS ENCODING THE SAME

<130> P3121R1

<140> US 09/964,994

<141> 2001-09-26

<150> PCT/US00/08439

<151> 2000-03-30

<150> PCT/US01/06520

<151> 2001-02-28

<150> US 60/191,015

<151> 2000-03-21

<150> US 09/941,992

<151> 2001-08-28

<160> 7

<210> 1

<211> 1318

<212> DNA

<213> Homo Sapien

<400> 1

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accaaaagga agacagcatc tgtttcctct ttggtcctga gctggttaaa 200
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 <211> 262
 <212> PRT
 <213> Homo Sapien

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 20 25 30
 Gln Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln
 35 40 45
 Trp Gln Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe
 50 55 60
 Val Gln Tyr Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His
 65 70 75
 Gln Lys Pro Ser Gly Cys Trp Gln His Ile Ser Cys Asn Phe Pro
 80 85 90
 Gly Cys Arg Thr Leu Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn
 95 100 105
 Lys Glu Asp Cys Trp Gly Thr Gln Glu Leu Ser Cys Asp Leu Thr
 110 115 120
 Ser Glu Thr Ser Asp Ile Gln Glu Pro Tyr Tyr Gly Arg Val Arg
 125 130 135
 Ala Ala Ser Ala Gly Ser Tyr Ser Glu Trp Ser Met Thr Pro Arg
 140 145 150
 Phe Thr Pro Trp Trp Glu Thr Lys Ile Asp Pro Pro Val Met Asn
 155 160 165

Ile	Thr	Gln	Val	Asn	Gly	Ser	Leu	Leu	Val	Ile	Leu	His	Ala	Pro
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Asn	Leu	Pro	Tyr	Arg	Tyr	Gln	Lys	Glu	Lys	Asn	Val	Ser	Ile	Glu
				185					190					195
Asp	Tyr	Tyr	Glu	Leu	Leu	Tyr	Arg	Val	Phe	Ile	Ile	Asn	Asn	Ser
				200					205					210
Leu	Glu	Lys	Glu	Gln	Lys	Val	Tyr	Glu	Gly	Ala	His	Arg	Ala	Val
				215					220					225
Glu	Ile	Glu	Ala	Leu	Thr	Pro	His	Ser	Ser	Tyr	Cys	Val	Val	Ala
				230					235					240
Glu	Ile	Tyr	Gln	Pro	Met	Leu	Asp	Arg	Arg	Ser	Gln	Arg	Ser	Glu
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 <213> Artificial Sequence

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 <223> Synthetic oligonucleotide probe

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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 4
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<210> 5
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<220>
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gaagaattga ggctgcttgg gaggaaggcc aggaggaaca cgagactgag 250
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cagaaactgt ggaagcctt ctgggctgtg aaagacacta tgcaagctca 500
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gaaggtgcct ctggatgctg tgaagagtct acagagaaga ttcttgtatt 1100
tattacaact ctatttaatt aatgtcagta tttcaactga agttctattt 1150
atttgtgaga ctgtaagtta catgaaggca gcagaatatt gtgccccatg 1200
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